|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Gene\_id** | **Rename** | **Description** | **FPKM** | | |
| **CK** | **MD\_2h** | **MD\_6h** |
| TRINITY\_DN45628\_c2\_g11 | CcMYB1 | Myb-like DNA-binding domain | 0.15 | 0.71 | 0.65 |
| TRINITY\_DN44798\_c2\_g4 | CcMYB2 | Myb-like DNA-binding domain | 0.00 | 25.39 | 6.22 |
| TRINITY\_DN49151\_c1\_g1 | CcMYB3 | Myb-like DNA-binding domain | 17.21 | 18.20 | 3.16 |
| TRINITY\_DN50766\_c3\_g1 | CcMYB4 | Myb-like DNA-binding domain | 1.73 | 0.22 | 2.77 |
| TRINITY\_DN31698\_c0\_g1 | CcMYB5 | Myb-like DNA-binding domain | 29.63 | 11.38 | 3.28 |
| TRINITY\_DN40236\_c0\_g1 | CcMYB6 | Myb-like DNA-binding domain | 42.64 | 6.38 | 9.81 |
| TRINITY\_DN50575\_c2\_g1 | CcMYB7 | Myb-like DNA-binding domain | 5.60 | 1.30 | 8.34 |
| TRINITY\_DN42715\_c0\_g3 | CcMYB8 | Myb-like DNA-binding domain | 7.91 | 138.32 | 79.28 |
| TRINITY\_DN42715\_c1\_g1 | CcMYB9 | Myb-like DNA-binding domain | 7.10 | 3.65 | 5.53 |
| TRINITY\_DN49154\_c3\_g1 | CcMYB10 | Myb-like DNA-binding domain | 0.00 | 0.66 | 1.09 |
| TRINITY\_DN49195\_c1\_g1 | CcMYB11 | Myb-like DNA-binding domain | 11.30 | 6.51 | 5.06 |
| TRINITY\_DN49195\_c2\_g2 | CcMYB12 | Myb-like DNA-binding domain | 18.24 | 7.99 | 9.97 |
| TRINITY\_DN47421\_c0\_g1 | CcMYB13 | Myb-like DNA-binding domain | 6.70 | 2.08 | 3.24 |
| TRINITY\_DN48475\_c1\_g1 | CcMYB14 | Myb-like DNA-binding domain | 0.03 | 0.68 | 0.78 |
| TRINITY\_DN50385\_c2\_g1 | CcMYB15 | Myb-like DNA-binding domain | 0.05 | 0.75 | 1.00 |
| TRINITY\_DN36467\_c0\_g1 | CcMYB16 | Myb-like DNA-binding domain | 3.26 | 3.44 | 1.88 |
| TRINITY\_DN45791\_c3\_g2 | CcMYB17 | Myb-like DNA-binding domain | 40.35 | 26.96 | 65.27 |
| TRINITY\_DN46797\_c4\_g1 | CcMYB18 | Myb-like DNA-binding domain | 2.11 | 2.74 | 6.03 |
| TRINITY\_DN46797\_c4\_g3 | CcMYB19 | Myb-like DNA-binding domain | 2.66 | 1.59 | 6.88 |
| TRINITY\_DN51230\_c1\_g1 | CcMYB20 | Myb-like DNA-binding domain | 31.43 | 12.94 | 12.89 |
| TRINITY\_DN51230\_c1\_g8 | CcMYB21 | Myb-like DNA-binding domain | 0.45 | 0.09 | 1.42 |
| TRINITY\_DN48168\_c0\_g1 | CcMYB22 | Myb-like DNA-binding domain | 6.66 | 6.72 | 3.68 |
| TRINITY\_DN48168\_c1\_g1 | CcMYB23 | Myb-like DNA-binding domain | 6.38 | 7.01 | 27.57 |
| TRINITY\_DN48168\_c4\_g2 | CcMYB24 | Myb-like DNA-binding domain | 6.34 | 5.66 | 95.65 |
| TRINITY\_DN43012\_c0\_g1 | CcMYB25 | Myb-like DNA-binding domain | 0.05 | 0.19 | 1.83 |
| TRINITY\_DN45267\_c2\_g3 | CcMYB26 | Myb-like DNA-binding domain | 0.69 | 25.60 | 12.74 |
| TRINITY\_DN51412\_c3\_g1 | CcMYB27 | Myb-like DNA-binding domain | 15.25 | 30.64 | 6.46 |
| TRINITY\_DN43639\_c1\_g1 | CcMYB28 | Myb-like DNA-binding domain | 39.32 | 20.74 | 20.89 |
| TRINITY\_DN43639\_c1\_g6 | CcMYB29 | Myb-like DNA-binding domain | 3.06 | 6.96 | 0.67 |
| TRINITY\_DN43639\_c1\_g5 | CcMYB30 | Myb-like DNA-binding domain | 0.71 | 0.25 | 0.18 |
| TRINITY\_DN41214\_c1\_g2 | CcMYB31 | Myb-like DNA-binding domain | 5.85 | 2.06 | 3.72 |
| TRINITY\_DN41272\_c0\_g1 | CcMYB32 | Myb-like DNA-binding domain | 1.42 | 41.37 | 37.35 |
| TRINITY\_DN25581\_c0\_g1 | CcMYB33 | Myb-like DNA-binding domain | 1.31 | 2.18 | 0.71 |
| TRINITY\_DN43044\_c0\_g2 | CcMYB34 | Myb-like DNA-binding domain | 0.35 | 2.27 | 0.93 |
| TRINITY\_DN44712\_c1\_g1 | CcMYB35 | Myb-like DNA-binding domain | 2.09 | 2.00 | 0.31 |
| TRINITY\_DN44713\_c0\_g1 | CcMYB36 | Myb-like DNA-binding domain | 268.83 | 269.04 | 93.44 |
| TRINITY\_DN44707\_c0\_g1 | CcMYB37 | Myb-like DNA-binding domain | 39.39 | 32.08 | 21.13 |
| TRINITY\_DN44728\_c7\_g1 | CcMYB38 | Myb-like DNA-binding domain | 0.20 | 0.12 | 1.00 |
| TRINITY\_DN47529\_c4\_g2 | CcMYB39 | Myb-like DNA-binding domain | 75.67 | 51.44 | 29.41 |
| TRINITY\_DN49235\_c1\_g1 | CcMYB40 | Myb-like DNA-binding domain | 0.36 | 1.11 | 7.46 |
| TRINITY\_DN49627\_c1\_g2 | CcMYB41 | Myb-like DNA-binding domain | 3.07 | 18.72 | 12.32 |
| TRINITY\_DN50844\_c0\_g2 | CcMYB42 | Myb-like DNA-binding domain | 46.35 | 51.74 | 25.52 |
| TRINITY\_DN46101\_c2\_g7 | CcMYB43 | Myb-like DNA-binding domain | 40.50 | 6.68 | 34.04 |
| TRINITY\_DN41900\_c0\_g1 | CcMYB44 | Myb-like DNA-binding domain | 2.13 | 5.56 | 3.68 |
| TRINITY\_DN48422\_c1\_g4 | CcMYB45 | Myb-like DNA-binding domain | 7.39 | 4.52 | 3.87 |
| TRINITY\_DN50238\_c2\_g2 | CcMYB46 | Myb-like DNA-binding domain | 11.53 | 19.77 | 13.40 |
| TRINITY\_DN46238\_c1\_g1 | CcMYB47 | Myb-like DNA-binding domain | 26.64 | 26.01 | 8.72 |
| TRINITY\_DN39982\_c0\_g1 | CcMYB48 | Myb-like DNA-binding domain | 3.54 | 2.01 | 3.38 |
| TRINITY\_DN43622\_c2\_g1 | CcMYB49 | Myb-like DNA-binding domain | 0.00 | 0.57 | 0.70 |
| TRINITY\_DN43663\_c0\_g1 | CcMYB50 | Myb-like DNA-binding domain | 13.64 | 14.80 | 74.34 |
| TRINITY\_DN43571\_c1\_g5 | CcMYB51 | Myb-like DNA-binding domain | 1.09 | 1.62 | 4.53 |
| TRINITY\_DN37951\_c0\_g1 | CcMYB52 | Myb-like DNA-binding domain | 0.17 | 1.96 | 6.48 |
| TRINITY\_DN47044\_c1\_g1 | CcMYB53 | Myb-like DNA-binding domain | 0.05 | 0.38 | 1.27 |
| TRINITY\_DN50733\_c0\_g2 | CcMYB54 | Myb-like DNA-binding domain | 0.89 | 12.46 | 30.56 |
| TRINITY\_DN43903\_c2\_g1 | CcMYB55 | Myb-like DNA-binding domain | 111.86 | 76.58 | 24.47 |
| TRINITY\_DN46417\_c6\_g1 | CcMYB56 | Myb-like DNA-binding domain | 2.40 | 7.66 | 16.43 |
| TRINITY\_DN44150\_c0\_g3 | CcMYB57 | Myb-like DNA-binding domain | 3.32 | 2.46 | 5.50 |
| TRINITY\_DN47613\_c0\_g1 | CcMYB58 | Myb-like DNA-binding domain | 1.52 | 2.35 | 0.95 |
| TRINITY\_DN46626\_c1\_g1 | CcMYB59 | Myb-like DNA-binding domain | 3.70 | 7.79 | 3.08 |
| TRINITY\_DN46626\_c1\_g1 | CcMYB60 | Myb-like DNA-binding domain | 19.34 | 5.38 | 19.16 |
| TRINITY\_DN44653\_c5\_g2 | CcMYB61 | Myb-like DNA-binding domain | 75.09 | 15.81 | 24.62 |
| TRINITY\_DN42859\_c3\_g4 | CcMYB62 | Myb-like DNA-binding domain | 345.38 | 109.39 | 94.71 |
| TRINITY\_DN42859\_c3\_g1 | CcMYB63 | Myb-like DNA-binding domain | 15.37 | 62.49 | 97.72 |
| TRINITY\_DN48132\_c1\_g2 | CcMYB64 | Myb-like DNA-binding domain | 2.51 | 4.74 | 1.78 |
| TRINITY\_DN47090\_c1\_g1 | CcMYB65 | Myb-like DNA-binding domain | 61.63 | 20.26 | 24.29 |
| TRINITY\_DN47090\_c1\_g1 | CcMYB66 | Myb-like DNA-binding domain | 8.76 | 2.81 | 23.78 |
| TRINITY\_DN47090\_c1\_g2 | CcMYB67 | Myb-like DNA-binding domain | 0.33 | 0.21 | 2.05 |
| TRINITY\_DN47090\_c1\_g4 | CcMYB68 | Myb-like DNA-binding domain | 0.10 | 108.87 | 24.82 |
| TRINITY\_DN47090\_c1\_g3 | CcMYB69 | Myb-like DNA-binding domain | 3.55 | 7.72 | 13.65 |
| TRINITY\_DN31376\_c0\_g1 | CcMYB70 | Myb-like DNA-binding domain | 46.77 | 21.56 | 29.95 |
| TRINITY\_DN49768\_c1\_g4 | CcMYB71 | Myb-like DNA-binding domain | 14.69 | 12.93 | 8.83 |
| TRINITY\_DN49787\_c0\_g1 | CcMYB72 | Myb-like DNA-binding domain | 12.16 | 13.62 | 5.09 |
| TRINITY\_DN43465\_c2\_g3 | CcMYB73 | Myb-like DNA-binding domain | 7.85 | 6.24 | 0.81 |
| TRINITY\_DN43465\_c2\_g1 | CcMYB74 | Myb-like DNA-binding domain | 9.34 | 6.79 | 2.19 |
| TRINITY\_DN47910\_c1\_g2 | CcMYB75 | Myb-like DNA-binding domain | 192.87 | 67.31 | 43.87 |
| TRINITY\_DN46359\_c3\_g6 | CcMYB76 | Myb-like DNA-binding domain | 24.03 | 5.50 | 3.53 |
| TRINITY\_DN40317\_c0\_g1 | CcMYB77 | Myb-like DNA-binding domain | 0.05 | 1.83 | 2.52 |
| TRINITY\_DN43933\_c1\_g1 | CcMYB78 | Myb-like DNA-binding domain | 120.92 | 22.97 | 56.86 |
| TRINITY\_DN44036\_c1\_g6 | CcMYB79 | Myb-like DNA-binding domain | 10.53 | 3.71 | 6.34 |
| TRINITY\_DN49926\_c0\_g1 | CcMYB80 | Myb-like DNA-binding domain | 9.58 | 19.42 | 5.05 |
| TRINITY\_DN36663\_c0\_g2 | CcMYB81 | Myb-like DNA-binding domain | 0.35 | 12.31 | 47.34 |
| TRINITY\_DN42710\_c1\_g2 | CcMYB82 | Myb-like DNA-binding domain | 10.13 | 44.97 | 16.44 |
| TRINITY\_DN43422\_c1\_g9 | CcMYB83 | Myb-like DNA-binding domain | 0.77 | 15.86 | 4.26 |
| TRINITY\_DN47700\_c0\_g1 | CcMYB84 | Myb-like DNA-binding domain | 0.61 | 24.82 | 88.55 |
| TRINITY\_DN47700\_c1\_g4 | CcMYB85 | Myb-like DNA-binding domain | 0.55 | 3.00 | 30.16 |
| TRINITY\_DN47700\_c1\_g6 | CcMYB86 | Myb-like DNA-binding domain | 0.38 | 4.94 | 5.47 |
| TRINITY\_DN47700\_c1\_g1 | CcMYB87 | Myb-like DNA-binding domain | 0.72 | 2.35 | 2.51 |
| TRINITY\_DN44045\_c1\_g7 | CcMYB88 | Myb-like DNA-binding domain | 24.62 | 105.46 | 34.87 |
| TRINITY\_DN48393\_c2\_g1 | CcMYB89 | Myb-like DNA-binding domain | 2.23 | 2.58 | 9.89 |
| TRINITY\_DN48393\_c3\_g4 | CcMYB90 | Myb-like DNA-binding domain | 1.01 | 3.22 | 2.25 |
| TRINITY\_DN48393\_c3\_g2 | CcMYB91 | Myb-like DNA-binding domain | 3.80 | 0.60 | 7.82 |
| TRINITY\_DN50435\_c0\_g1 | CcMYB92 | Myb-like DNA-binding domain | 0.83 | 3.53 | 19.99 |
| TRINITY\_DN44375\_c1\_g3 | CcMYB93 | Myb-like DNA-binding domain | 0.34 | 1.95 | 3.12 |
| TRINITY\_DN37633\_c0\_g1 | CcMYB94 | Myb-like DNA-binding domain | 0.88 | 129.47 | 35.34 |
| TRINITY\_DN48353\_c2\_g3 | CcMYB95 | Myb-like DNA-binding domain | 0.05 | 5.03 | 0.57 |
| TRINITY\_DN48353\_c2\_g4 | CcMYB96 | Myb-like DNA-binding domain | 3.40 | 45.36 | 22.60 |
| TRINITY\_DN45916\_c5\_g1 | CcMYB97 | Myb-like DNA-binding domain | 0.36 | 5.38 | 2.74 |
| TRINITY\_DN42397\_c0\_g1 | CcMYB98 | Myb-like DNA-binding domain | 94.90 | 145.53 | 75.03 |
| TRINITY\_DN48049\_c1\_g1 | CcMYB99 | Myb-like DNA-binding domain | 10.96 | 22.75 | 30.62 |
| TRINITY\_DN48054\_c1\_g1 | CcMYB100 | Myb-like DNA-binding domain | 0.59 | 0.35 | 1.86 |
| TRINITY\_DN48054\_c2\_g2 | CcMYB101 | Myb-like DNA-binding domain | 12.79 | 6.28 | 4.05 |
| TRINITY\_DN48054\_c2\_g3 | CcMYB102 | Myb-like DNA-binding domain | 5.66 | 5.12 | 1.90 |
| TRINITY\_DN48054\_c2\_g7 | CcMYB103 | Myb-like DNA-binding domain | 0.36 | 55.92 | 17.08 |
| TRINITY\_DN48054\_c2\_g8 | CcMYB104 | Myb-like DNA-binding domain | 16.79 | 6.35 | 14.42 |
| TRINITY\_DN47691\_c1\_g6 | CcMYB105 | Myb-like DNA-binding domain | 18.18 | 19.49 | 8.81 |
| TRINITY\_DN47691\_c1\_g5 | CcMYB106 | Myb-like DNA-binding domain | 2.83 | 6.97 | 5.77 |
| TRINITY\_DN50985\_c1\_g4 | CcMYB107 | Myb-like DNA-binding domain | 37.60 | 21.92 | 58.73 |
| TRINITY\_DN49375\_c0\_g5 | CcMYB108 | Myb-like DNA-binding domain | 0.48 | 1.53 | 0.21 |
| TRINITY\_DN41558\_c0\_g1 | CcMYB109 | Myb-like DNA-binding domain | 3.74 | 2.49 | 1.18 |
| TRINITY\_DN45409\_c0\_g1 | CcMYB110 | Myb-like DNA-binding domain | 50.45 | 145.89 | 292.68 |
| TRINITY\_DN45409\_c0\_g2 | CcMYB111 | Myb-like DNA-binding domain | 4.14 | 3.40 | 2.61 |
| TRINITY\_DN45409\_c0\_g9 | CcMYB112 | Myb-like DNA-binding domain | 6.09 | 4.37 | 0.50 |
| TRINITY\_DN45451\_c2\_g1 | CcMYB113 | Myb-like DNA-binding domain | 71.44 | 21.51 | 30.21 |
| TRINITY\_DN50794\_c1\_g1 | CcMYB114 | Myb-like DNA-binding domain | 304.71 | 243.80 | 45.60 |
| TRINITY\_DN34173\_c0\_g1 | CcMYB115 | Myb-like DNA-binding domain | 9.41 | 23.36 | 12.09 |
| TRINITY\_DN44521\_c0\_g1 | CcMYB116 | Myb-like DNA-binding domain | 35.19 | 80.23 | 61.72 |
| TRINITY\_DN38902\_c0\_g1 | CcMYB117 | Myb-like DNA-binding domain | 6.25 | 19.95 | 24.40 |
| TRINITY\_DN51074\_c8\_g1 | CcMYB118 | Myb-like DNA-binding domain | 0.03 | 0.38 | 0.98 |
| TRINITY\_DN42970\_c0\_g1 | CcMYB119 | Myb-like DNA-binding domain | 54.81 | 18.23 | 28.26 |
| TRINITY\_DN44966\_c2\_g4 | CcMYB120 | Myb-like DNA-binding domain | 3.28 | 70.42 | 70.35 |
| TRINITY\_DN48238\_c0\_g1 | CcMYB121 | Myb-like DNA-binding domain | 1.32 | 3.45 | 0.89 |
| TRINITY\_DN48238\_c0\_g3 | CcMYB122 | Myb-like DNA-binding domain | 1.21 | 4.13 | 0.70 |
| TRINITY\_DN48227\_c3\_g1 | CcMYB123 | Myb-like DNA-binding domain | 66.49 | 7.48 | 48.73 |
| TRINITY\_DN48227\_c3\_g5 | CcMYB124 | Myb-like DNA-binding domain | 17.98 | 7.10 | 12.28 |
| TRINITY\_DN45569\_c2\_g1 | CcMYB125 | Myb-like DNA-binding domain | 4.60 | 2.88 | 0.80 |
| TRINITY\_DN19910\_c0\_g1 | CcMYB126 | Myb-like DNA-binding domain | 6.25 | 19.95 | 24.40 |
| TRINITY\_DN40431\_c0\_g1 | CcMYB127 | Myb-like DNA-binding domain | 0.03 | 0.38 | 0.98 |
| TRINITY\_DN35704\_c0\_g1 | CcWRKY1 | WRKY DNA -binding domain | 4.56 | 1.13 | 3.34 |
| TRINITY\_DN45791\_c4\_g1 | CcWRKY2 | WRKY DNA -binding domain | 0.18 | 0.67 | 0.73 |
| TRINITY\_DN46744\_c1\_g5 | CcWRKY3 | WRKY DNA -binding domain | 0.76 | 5.12 | 2.85 |
| TRINITY\_DN50241\_c3\_g3 | CcWRKY4 | WRKY DNA -binding domain | 1.91 | 1.89 | 0.53 |
| TRINITY\_DN51409\_c2\_g3 | CcWRKY5 | WRKY DNA -binding domain | 0.03 | 0.30 | 3.82 |
| TRINITY\_DN51409\_c2\_g9 | CcWRKY6 | WRKY DNA -binding domain | 2.40 | 6.15 | 3.71 |
| TRINITY\_DN45494\_c1\_g4 | CcWRKY7 | WRKY DNA -binding domain | 0.31 | 2.96 | 1.93 |
| TRINITY\_DN48629\_c0\_g1 | CcWRKY8 | WRKY DNA -binding domain | 108.09 | 927.99 | 128.19 |
| TRINITY\_DN48629\_c0\_g7 | CcWRKY9 | WRKY DNA -binding domain | 1.12 | 5.49 | 1.45 |
| TRINITY\_DN48228\_c0\_g10 | CcWRKY10 | WRKY DNA -binding domain | 0.51 | 0.26 | 7.09 |
| TRINITY\_DN49750\_c0\_g3 | CcWRKY11 | WRKY DNA -binding domain | 6.25 | 18.25 | 51.92 |
| TRINITY\_DN22541\_c0\_g2 | CcWRKY12 | WRKY DNA -binding domain | 0.04 | 6.05 | 1.12 |
| TRINITY\_DN46259\_c0\_g4 | CcWRKY13 | WRKY DNA -binding domain | 1.96 | 32.14 | 11.11 |
| TRINITY\_DN46587\_c2\_g1 | CcWRKY14 | WRKY DNA -binding domain | 9.18 | 35.00 | 57.59 |
| TRINITY\_DN46572\_c2\_g1 | CcWRKY15 | WRKY DNA -binding domain | 9.16 | 8.08 | 26.18 |
| TRINITY\_DN38622\_c0\_g1 | CcWRKY16 | WRKY DNA -binding domain | 7.32 | 2.43 | 3.74 |
| TRINITY\_DN49545\_c1\_g1 | CcWRKY17 | WRKY DNA -binding domain | 8.00 | 322.41 | 137.58 |
| TRINITY\_DN44317\_c2\_g1 | CcWRKY18 | WRKY DNA -binding domain | 6.36 | 15.36 | 17.62 |
| TRINITY\_DN48296\_c3\_g1 | CcWRKY19 | WRKY DNA -binding domain | 1.07 | 0.48 | 0.54 |
| TRINITY\_DN39627\_c0\_g2 | CcWRKY20 | WRKY DNA -binding domain | 37.85 | 10.67 | 10.86 |
| TRINITY\_DN38119\_c0\_g1 | CcWRKY21 | WRKY DNA -binding domain | 2.40 | 37.37 | 118.20 |
| TRINITY\_DN45810\_c3\_g4 | CcWRKY22 | WRKY DNA -binding domain | 25.96 | 293.60 | 222.89 |
| TRINITY\_DN48306\_c1\_g3 | CcWRKY23 | WRKY DNA -binding domain | 1.18 | 219.66 | 135.36 |
| TRINITY\_DN43541\_c3\_g1 | CcWRKY24 | WRKY DNA -binding domain | 59.45 | 232.01 | 109.07 |
| TRINITY\_DN43541\_c3\_g2 | CcWRKY25 | WRKY DNA -binding domain | 2.35 | 1.74 | 8.81 |
| TRINITY\_DN43541\_c3\_g3 | CcWRKY26 | WRKY DNA -binding domain | 3.01 | 55.75 | 54.98 |
| TRINITY\_DN46028\_c0\_g1 | CcWRKY27 | WRKY DNA -binding domain | 0.40 | 1.30 | 0.69 |
| TRINITY\_DN42888\_c0\_g1 | CcWRKY28 | WRKY DNA -binding domain | 5.09 | 7.45 | 36.99 |
| TRINITY\_DN46617\_c0\_g1 | CcWRKY29 | WRKY DNA -binding domain | 4.96 | 3.75 | 8.58 |
| TRINITY\_DN48000\_c1\_g1 | CcWRKY30 | WRKY DNA -binding domain | 39.25 | 306.36 | 215.00 |
| TRINITY\_DN48000\_c1\_g1 | CcWRKY31 | WRKY DNA -binding domain | 7.12 | 91.05 | 38.62 |
| TRINITY\_DN43812\_c0\_g7 | CcWRKY32 | WRKY DNA -binding domain | 24.58 | 79.33 | 25.41 |
| TRINITY\_DN45566\_c3\_g1 | CcWRKY33 | WRKY DNA -binding domain | 37.45 | 625.83 | 225.35 |
| TRINITY\_DN44923\_c1\_g2 | CcWRKY34 | WRKY DNA -binding domain | 0.33 | 0.08 | 2.47 |
| TRINITY\_DN44923\_c1\_g7 | CcWRKY35 | WRKY DNA -binding domain | 3.14 | 6.42 | 0.26 |
| TRINITY\_DN44923\_c1\_g8 | CcWRKY36 | WRKY DNA -binding domain | 0.63 | 66.87 | 22.79 |
| TRINITY\_DN44923\_c1\_g1 | CcWRKY37 | WRKY DNA -binding domain | 7.12 | 91.05 | 38.62 |
| TRINITY\_DN49652\_c0\_g5 | CcbZIP1 | bZIP transcription factor | 0.15 | 0.13 | 0.85 |
| TRINITY\_DN44798\_c4\_g2 | CcbZIP2 | bZIP transcription factor | 2.06 | 0.47 | 0.66 |
| TRINITY\_DN40264\_c2\_g2 | CcbZIP3 | bZIP transcription factor | 0.42 | 0.04 | 0.65 |
| TRINITY\_DN44382\_c0\_g1 | CcbZIP4 | bZIP transcription factor | 0.14 | 0.11 | 0.94 |
| TRINITY\_DN36888\_c0\_g2 | CcbZIP5 | bZIP transcription factor | 0.00 | 0.21 | 1.11 |
| TRINITY\_DN50241\_c0\_g1 | CcbZIP6 | bZIP transcription factor | 0.13 | 0.19 | 2.56 |
| TRINITY\_DN51472\_c2\_g2 | CcbZIP7 | bZIP transcription factor | 0.31 | 0.12 | 0.83 |
| TRINITY\_DN51472\_c2\_g6 | CcbZIP8 | bZIP transcription factor | 0.12 | 0.10 | 0.79 |
| TRINITY\_DN43628\_c0\_g3 | CcbZIP9 | bZIP transcription factor | 4.62 | 1.80 | 2.60 |
| TRINITY\_DN41214\_c1\_g1 | CcbZIP10 | bZIP transcription factor | 3.30 | 2.78 | 7.28 |
| TRINITY\_DN47722\_c0\_g5 | CcbZIP11 | bZIP transcription factor | 12.35 | 19.16 | 7.74 |
| TRINITY\_DN44701\_c0\_g1 | CcbZIP12 | bZIP transcription factor | 0.00 | 1.16 | 1.42 |
| TRINITY\_DN46543\_c0\_g1 | CcbZIP13 | bZIP transcription factor | 1.85 | 31.81 | 1.80 |
| TRINITY\_DN38629\_c1\_g1 | CcbZIP14 | bZIP transcription factor | 4.28 | 0.83 | 0.10 |
| TRINITY\_DN48835\_c1\_g1 | CcbZIP15 | bZIP transcription factor | 26.07 | 32.29 | 14.31 |
| TRINITY\_DN20038\_c0\_g1 | CcbZIP16 | bZIP transcription factor | 0.02 | 2.69 | 1.21 |
| TRINITY\_DN47036\_c3\_g1 | CcbZIP17 | bZIP transcription factor | 3.80 | 24.97 | 9.23 |
| TRINITY\_DN47036\_c3\_g2 | CcbZIP18 | bZIP transcription factor | 0.05 | 0.18 | 1.86 |
| TRINITY\_DN51237\_c1\_g3 | CcbZIP19 | bZIP transcription factor | 0.05 | 3.84 | 2.85 |
| TRINITY\_DN27565\_c0\_g2 | CcbZIP20 | bZIP transcription factor | 0.30 | 2.32 | 3.93 |
| TRINITY\_DN38123\_c0\_g1 | CcbZIP21 | bZIP transcription factor | 1.31 | 7.96 | 11.11 |
| TRINITY\_DN35256\_c0\_g1 | CcbZIP22 | bZIP transcription factor | 0.08 | 5.77 | 4.27 |
| TRINITY\_DN48070\_c0\_g1 | CcbZIP23 | bZIP transcription factor | 0.04 | 1.76 | 1.68 |
| TRINITY\_DN43569\_c0\_g1 | CcbZIP24 | bZIP transcription factor | 0.06 | 0.42 | 1.25 |
| TRINITY\_DN44375\_c1\_g4 | CcbZIP25 | bZIP transcription factor | 1.87 | 0.44 | 2.32 |
| TRINITY\_DN46043\_c0\_g1 | CcbZIP26 | bZIP transcription factor | 17.87 | 6.56 | 15.42 |
| TRINITY\_DN50994\_c6\_g5 | CcbZIP27 | bZIP transcription factor | 29.63 | 147.90 | 74.92 |
| TRINITY\_DN6958\_c0\_g1 | CcbZIP28 | bZIP transcription factor | 4.48 | 5.72 | 37.29 |
| TRINITY\_DN48868\_c0\_g3 | CcbZIP29 | bZIP transcription factor | 0.96 | 10.81 | 2.59 |
| TRINITY\_DN41548\_c0\_g1 | CcbZIP30 | bZIP transcription factor | 4.86 | 4.97 | 2.98 |
| TRINITY\_DN37197\_c0\_g2 | CcbZIP31 | bZIP transcription factor | 2.14 | 0.48 | 0.78 |
| TRINITY\_DN43356\_c0\_g1 | CcbZIP32 | bZIP transcription factor | 1.16 | 0.31 | 1.04 |
| TRINITY\_DN45559\_c0\_g1 | CcbZIP33 | bZIP transcription factor | 1.67 | 5.28 | 10.26 |
| TRINITY\_DN44555\_c3\_g1 | CcbZIP34 | bZIP transcription factor | 4.35 | 4.63 | 10.46 |
| TRINITY\_DN46333\_c2\_g1 | CcbZIP35 | bZIP transcription factor | 0.64 | 1.28 | 0.32 |
| TRINITY\_DN44233\_c0\_g1 | CcERF1 | AP2 domain | 0.00 | 1.08 | 0.56 |
| TRINITY\_DN44036\_c1\_g2 | CcERF2 | AP2 domain | 0.75 | 0.72 | 0.17 |
| TRINITY\_DN14404\_c0\_g1 | CcERF3 | AP2 domain | 0.06 | 0.73 | 0.78 |
| TRINITY\_DN45121\_c2\_g3 | CcERF4 | AP2 domain | 8.46 | 30.40 | 25.00 |
| TRINITY\_DN46170\_c1\_g1 | CcERF5 | AP2 domain | 0.13 | 131.89 | 58.94 |
| TRINITY\_DN46170\_c1\_g2 | CcERF6 | AP2 domain | 1.39 | 30.87 | 29.64 |
| TRINITY\_DN46170\_c1\_g3 | CcERF7 | AP2 domain | 0.00 | 0.25 | 1.23 |
| TRINITY\_DN46170\_c1\_g4 | CcERF8 | AP2 domain | 0.65 | 2.66 | 5.18 |
| TRINITY\_DN46170\_c1\_g5 | CcERF9 | AP2 domain | 0.02 | 14.95 | 9.53 |
| TRINITY\_DN46170\_c1\_g6 | CcERF10 | AP2 domain | 0.00 | 73.99 | 18.29 |
| TRINITY\_DN46170\_c1\_g7 | CcERF11 | AP2 domain | 4.31 | 15.99 | 15.16 |
| TRINITY\_DN46170\_c1\_g8 | CcERF12 | AP2 domain | 32.03 | 138.82 | 103.33 |
| TRINITY\_DN41203\_c0\_g2 | CcERF13 | AP2 domain | 0.35 | 169.46 | 66.90 |
| TRINITY\_DN49918\_c2\_g2 | CcERF14 | AP2 domain | 0.19 | 131.25 | 72.43 |
| TRINITY\_DN49918\_c3\_g1 | CcERF15 | AP2 domain | 0.00 | 6.25 | 2.87 |
| TRINITY\_DN49246\_c0\_g3 | CcERF16 | AP2 domain | 0.15 | 2.33 | 1.55 |
| TRINITY\_DN49246\_c0\_g1 | CcERF17 | AP2 domain | 2.22 | 3.94 | 5.65 |
| TRINITY\_DN49246\_c0\_g4 | CcERF18 | AP2 domain | 0.82 | 41.21 | 159.80 |
| TRINITY\_DN36480\_c0\_g1 | CcERF19 | AP2 domain | 0.88 | 7.03 | 7.58 |
| TRINITY\_DN51227\_c0\_g3 | CcERF20 | AP2 domain | 0.02 | 125.71 | 79.02 |
| TRINITY\_DN62408\_c0\_g1 | CcERF21 | AP2 domain | 31.29 | 316.47 | 55.25 |
| TRINITY\_DN46791\_c3\_g5 | CcERF22 | AP2 domain | 65.09 | 588.47 | 505.10 |
| TRINITY\_DN43658\_c2\_g9 | CcERF23 | AP2 domain | 8.46 | 166.29 | 93.83 |
| TRINITY\_DN37494\_c0\_g1 | CcERF24 | AP2 domain | 3.82 | 253.61 | 117.33 |
| TRINITY\_DN45270\_c0\_g3 | CcERF25 | AP2 domain | 3.85 | 36.40 | 13.50 |
| TRINITY\_DN49492\_c2\_g7 | CcERF26 | AP2 domain | 1.22 | 313.08 | 73.03 |
| TRINITY\_DN47880\_c0\_g2 | CcERF27 | AP2 domain | 0.00 | 11.15 | 6.15 |
| TRINITY\_DN47880\_c0\_g3 | CcERF28 | AP2 domain | 0.02 | 9.45 | 2.10 |
| TRINITY\_DN41971\_c0\_g1 | CcERF29 | AP2 domain | 9.56 | 6.44 | 3.32 |
| TRINITY\_DN49708\_c0\_g12 | CcERF30 | AP2 domain | 1.78 | 0.57 | 0.48 |
| TRINITY\_DN38017\_c0\_g1 | CcERF31 | AP2 domain | 0.51 | 0.26 | 1.51 |
| TRINITY\_DN34248\_c0\_g1 | CcERF32 | AP2 domain | 0.13 | 49.60 | 16.71 |
| TRINITY\_DN46054\_c1\_g1 | CcERF33 | AP2 domain | 0.00 | 30.05 | 1.36 |
| TRINITY\_DN46575\_c2\_g1 | CcERF34 | AP2 domain | 0.00 | 6.79 | 0.33 |
| TRINITY\_DN46575\_c4\_g1 | CcERF35 | AP2 domain | 0.14 | 41.15 | 3.61 |
| TRINITY\_DN39424\_c2\_g1 | CcERF36 | AP2 domain | 0.00 | 17.86 | 4.11 |
| TRINITY\_DN38620\_c0\_g1 | CcERF37 | AP2 domain | 0.13 | 5.94 | 107.08 |
| TRINITY\_DN48853\_c1\_g3 | CcERF38 | AP2 domain | 3.27 | 68.26 | 84.29 |
| TRINITY\_DN49514\_c0\_g1 | CcERF39 | AP2 domain | 0.32 | 3.67 | 3.27 |
| TRINITY\_DN49514\_c1\_g1 | CcERF40 | AP2 domain | 2.16 | 3.42 | 5.90 |
| TRINITY\_DN42179\_c0\_g1 | CcERF41 | AP2 domain | 16.87 | 3.91 | 7.72 |
| TRINITY\_DN47289\_c0\_g1 | CcERF42 | AP2 domain | 6.47 | 1.40 | 13.22 |
| TRINITY\_DN44650\_c0\_g1 | CcERF43 | AP2 domain | 1.64 | 0.92 | 2.33 |
| TRINITY\_DN44650\_c1\_g1 | CcERF44 | AP2 domain | 0.93 | 6.27 | 9.29 |
| TRINITY\_DN44650\_c1\_g2 | CcERF45 | AP2 domain | 3.21 | 315.14 | 47.66 |
| TRINITY\_DN45626\_c0\_g1 | CcERF46 | AP2 domain | 0.41 | 15.50 | 37.53 |
| TRINITY\_DN51395\_c0\_g2 | CcERF47 | AP2 domain | 19.89 | 63.02 | 50.70 |
| TRINITY\_DN38620\_c0\_g2 | CcERF48 | AP2 domain | 1.62 | 24.52 | 19.87 |
| TRINITY\_DN42028\_c0\_g1 | CcERF49 | AP2 domain | 560.51 | 195.60 | 181.98 |
| TRINITY\_DN46449\_c2\_g7 | CcERF50 | AP2 domain | 1.34 | 5.89 | 21.54 |
| TRINITY\_DN45591\_c1\_g2 | CcERF51 | AP2 domain | 4.95 | 54.98 | 19.17 |
| TRINITY\_DN44628\_c0\_g2 | CcERF52 | AP2 domain | 34.09 | 281.37 | 108.66 |
| TRINITY\_DN48643\_c0\_g3 | CcERF53 | AP2 domain | 12.35 | 12.98 | 103.01 |
| TRINITY\_DN48918\_c0\_g9 | CcERF54 | AP2 domain | 0.21 | 12.92 | 14.29 |
| TRINITY\_DN44939\_c1\_g6 | CcERF55 | AP2 domain | 0.12 | 165.15 | 99.57 |
| TRINITY\_DN31476\_c0\_g1 | CcERF56 | AP2 domain | 1.10 | 25.07 | 65.46 |
| TRINITY\_DN43789\_c0\_g3 | CcERF57 | AP2 domain | 2.39 | 62.54 | 79.03 |
| TRINITY\_DN41307\_c0\_g1 | CcERF58 | AP2 domain | 42.36 | 17.35 | 15.91 |
| TRINITY\_DN48573\_c0\_g1 | CcERF59 | AP2 domain | 12.46 | 4.66 | 36.68 |
| TRINITY\_DN48667\_c1\_g1 | CcERF60 | AP2 domain | 0.04 | 2.52 | 0.32 |
| TRINITY\_DN44847\_c0\_g2 | CcERF61 | AP2 domain | 0.19 | 615.10 | 179.84 |
| TRINITY\_DN25477\_c0\_g1 | CcERF62 | AP2 domain | 4.52 | 15.66 | 5.25 |
| TRINITY\_DN47856\_c0\_g2 | CcERF63 | AP2 domain | 0.07 | 139.73 | 19.60 |
| TRINITY\_DN50063\_c1\_g3 | CcERF64 | AP2 domain | 0.22 | 28.04 | 12.52 |
| TRINITY\_DN44171\_c0\_g1 | CcERF65 | AP2 domain | 7.74 | 148.83 | 51.18 |
| TRINITY\_DN44171\_c1\_g1 | CcERF66 | AP2 domain | 3.49 | 7.57 | 3.86 |
| TRINITY\_DN44171\_c1\_g2 | CcERF67 | AP2 domain | 4.08 | 4.66 | 10.16 |
| TRINITY\_DN50995\_c1\_g1 | CcERF68 | AP2 domain | 135.94 | 92.68 | 74.03 |
| TRINITY\_DN50995\_c1\_g4 | CcERF69 | AP2 domain | 9.57 | 2.50 | 4.36 |
| TRINITY\_DN55146\_c0\_g1 | CcERF70 | AP2 domain | 2.96 | 2.96 | 1.47 |
| TRINITY\_DN50712\_c0\_g3 | CcERF71 | AP2 domain | 40.56 | 9.49 | 2.70 |
| TRINITY\_DN41575\_c0\_g1 | CcERF72 | AP2 domain | 1.76 | 5.19 | 1.90 |
| TRINITY\_DN45331\_c1\_g5 | CcERF73 | AP2 domain | 16.44 | 173.99 | 90.00 |
| TRINITY\_DN44586\_c0\_g3 | CcERF74 | AP2 domain | 0.12 | 44.80 | 17.00 |
| TRINITY\_DN44586\_c0\_g8 | CcERF75 | AP2 domain | 2.60 | 70.86 | 7.35 |
| TRINITY\_DN44586\_c0\_g7 | CcERF76 | AP2 domain | 0.16 | 60.05 | 6.61 |
| TRINITY\_DN44586\_c0\_g9 | CcERF77 | AP2 domain | 7.26 | 3.28 | 7.06 |
| TRINITY\_DN44586\_c0\_g4 | CcERF78 | AP2 domain | 7.65 | 6.34 | 3.09 |
| TRINITY\_DN44586\_c0\_g10 | CcERF79 | AP2 domain | 1.42 | 2.89 | 15.66 |
| TRINITY\_DN43260\_c3\_g5 | CcERF80 | AP2 domain | 0.00 | 0.40 | 2.03 |
| TRINITY\_DN44038\_c0\_g1 | CcBHLH1 | Helix-loop-helix DNA-binding domain | 0.24 | 0.19 | 1.75 |
| TRINITY\_DN38527\_c0\_g1 | CcBHLH2 | Helix-loop-helix DNA-binding domain | 0.44 | 0.07 | 0.94 |
| TRINITY\_DN19706\_c0\_g1 | CcBHLH3 | Helix-loop-helix DNA-binding domain | 0.02 | 0.39 | 0.63 |
| TRINITY\_DN42693\_c1\_g1 | CcBHLH4 | Helix-loop-helix DNA-binding domain | 0.11 | 0.71 | 0.55 |
| TRINITY\_DN36828\_c0\_g2 | CcBHLH5 | Helix-loop-helix DNA-binding domain | 1.05 | 0.25 | 0.11 |
| TRINITY\_DN41060\_c0\_g1 | CcBHLH6 | Helix-loop-helix DNA-binding domain | 0.38 | 3.40 | 1.45 |
| TRINITY\_DN45351\_c1\_g3 | CcBHLH7 | Helix-loop-helix DNA-binding domain | 2.15 | 0.90 | 1.23 |
| TRINITY\_DN45351\_c1\_g1 | CcBHLH8 | Helix-loop-helix DNA-binding domain | 0.00 | 0.67 | 1.14 |
| TRINITY\_DN45351\_c1\_g4 | CcBHLH9 | Helix-loop-helix DNA-binding domain | 3.27 | 0.97 | 14.86 |
| TRINITY\_DN45368\_c0\_g1 | CcBHLH10 | Helix-loop-helix DNA-binding domain | 0.02 | 0.89 | 1.06 |
| TRINITY\_DN49163\_c2\_g1 | CcBHLH11 | Helix-loop-helix DNA-binding domain | 16.75 | 7.33 | 3.99 |
| TRINITY\_DN49345\_c0\_g1 | CcBHLH12 | Helix-loop-helix DNA-binding domain | 0.00 | 0.74 | 0.71 |
| TRINITY\_DN41404\_c0\_g1 | CcBHLH13 | Helix-loop-helix DNA-binding domain | 0.00 | 1.42 | 0.82 |
| TRINITY\_DN43268\_c0\_g7 | CcBHLH14 | Helix-loop-helix DNA-binding domain | 20.67 | 15.35 | 38.33 |
| TRINITY\_DN44271\_c2\_g1 | CcBHLH15 | Helix-loop-helix DNA-binding domain | 41.29 | 17.56 | 22.85 |
| TRINITY\_DN45349\_c1\_g1 | CcBHLH16 | Helix-loop-helix DNA-binding domain | 0.00 | 2.85 | 3.41 |
| TRINITY\_DN48629\_c0\_g5 | CcBHLH17 | Helix-loop-helix DNA-binding domain | 0.09 | 2.52 | 0.75 |
| TRINITY\_DN43084\_c0\_g5 | CcBHLH18 | Helix-loop-helix DNA-binding domain | 98.62 | 73.83 | 58.03 |
| TRINITY\_DN31133\_c0\_g2 | CcBHLH19 | Helix-loop-helix DNA-binding domain | 5.93 | 3.28 | 2.89 |
| TRINITY\_DN46163\_c5\_g1 | CcBHLH20 | Helix-loop-helix DNA-binding domain | 0.55 | 2.59 | 0.92 |
| TRINITY\_DN46019\_c0\_g1 | CcBHLH21 | Helix-loop-helix DNA-binding domain | 4.72 | 3.48 | 2.10 |
| TRINITY\_DN39922\_c0\_g1 | CcBHLH22 | Helix-loop-helix DNA-binding domain | 6.60 | 1.62 | 8.71 |
| TRINITY\_DN49805\_c0\_g1 | CcBHLH23 | Helix-loop-helix DNA-binding domain | 21.69 | 61.14 | 54.84 |
| TRINITY\_DN43530\_c0\_g1 | CcBHLH24 | Helix-loop-helix DNA-binding domain | 1.59 | 22.60 | 8.55 |
| TRINITY\_DN49815\_c1\_g8 | CcBHLH25 | Helix-loop-helix DNA-binding domain | 20.55 | 20.82 | 11.62 |
| TRINITY\_DN51231\_c0\_g2 | CcBHLH26 | Helix-loop-helix DNA-binding domain | 6.38 | 12.76 | 7.00 |
| TRINITY\_DN48847\_c1\_g1 | CcBHLH27 | Helix-loop-helix DNA-binding domain | 0.01 | 1.21 | 4.31 |
| TRINITY\_DN48847\_c1\_g5 | CcBHLH28 | Helix-loop-helix DNA-binding domain | 0.38 | 6.13 | 9.51 |
| TRINITY\_DN47384\_c0\_g3 | CcBHLH29 | Helix-loop-helix DNA-binding domain | 0.16 | 0.73 | 0.12 |
| TRINITY\_DN46790\_c2\_g1 | CcBHLH30 | Helix-loop-helix DNA-binding domain | 4.53 | 3.29 | 2.10 |
| TRINITY\_DN48296\_c2\_g1 | CcBHLH31 | Helix-loop-helix DNA-binding domain | 12.76 | 7.21 | 7.70 |
| TRINITY\_DN48270\_c2\_g3 | CcBHLH32 | Helix-loop-helix DNA-binding domain | 9.71 | 29.62 | 12.76 |
| TRINITY\_DN28067\_c0\_g1 | CcBHLH33 | Helix-loop-helix DNA-binding domain | 5.48 | 108.39 | 48.68 |
| TRINITY\_DN30426\_c0\_g3 | CcBHLH34 | Helix-loop-helix DNA-binding domain | 6.77 | 1006.33 | 97.68 |
| TRINITY\_DN47927\_c0\_g1 | CcBHLH35 | Helix-loop-helix DNA-binding domain | 15.55 | 6.11 | 6.51 |
| TRINITY\_DN43370\_c0\_g1 | CcBHLH36 | Helix-loop-helix DNA-binding domain | 69.16 | 188.74 | 152.68 |
| TRINITY\_DN45803\_c1\_g6 | CcBHLH37 | Helix-loop-helix DNA-binding domain | 2.81 | 48.67 | 15.50 |
| TRINITY\_DN45803\_c1\_g1 | CcBHLH38 | Helix-loop-helix DNA-binding domain | 14.20 | 103.91 | 43.93 |
| TRINITY\_DN37799\_c0\_g2 | CcBHLH39 | Helix-loop-helix DNA-binding domain | 25.40 | 10.73 | 4.33 |
| TRINITY\_DN34917\_c0\_g1 | CcBHLH40 | Helix-loop-helix DNA-binding domain | 8.32 | 1.64 | 14.28 |
| TRINITY\_DN37053\_c0\_g1 | CcBHLH41 | Helix-loop-helix DNA-binding domain | 23.18 | 211.67 | 36.75 |
| TRINITY\_DN28254\_c0\_g2 | CcBHLH42 | Helix-loop-helix DNA-binding domain | 0.41 | 83.36 | 12.08 |
| TRINITY\_DN42728\_c0\_g1 | CcBHLH43 | Helix-loop-helix DNA-binding domain | 254.55 | 149.46 | 95.77 |
| TRINITY\_DN44014\_c0\_g1 | CcBHLH44 | Helix-loop-helix DNA-binding domain | 5.43 | 3.33 | 1.49 |
| TRINITY\_DN45579\_c0\_g3 | CcBHLH45 | Helix-loop-helix DNA-binding domain | 1.02 | 243.99 | 157.07 |
| TRINITY\_DN46569\_c0\_g1 | CcBHLH46 | Helix-loop-helix DNA-binding domain | 0.34 | 0.24 | 1.08 |
| TRINITY\_DN48331\_c2\_g5 | CcBHLH47 | Helix-loop-helix DNA-binding domain | 24.22 | 104.46 | 42.15 |
| TRINITY\_DN43800\_c0\_g1 | CcBHLH48 | Helix-loop-helix DNA-binding domain | 0.82 | 52.56 | 6.68 |
| TRINITY\_DN47882\_c0\_g1 | CcBHLH49 | Helix-loop-helix DNA-binding domain | 97.62 | 59.02 | 20.03 |
| TRINITY\_DN49353\_c1\_g4 | CcBHLH50 | Helix-loop-helix DNA-binding domain | 2.64 | 10.56 | 3.99 |
| TRINITY\_DN45412\_c2\_g4 | CcBHLH51 | Helix-loop-helix DNA-binding domain | 17.51 | 15.62 | 7.46 |
| TRINITY\_DN50724\_c2\_g2 | CcBHLH52 | Helix-loop-helix DNA-binding domain | 5.23 | 8.01 | 3.97 |
| TRINITY\_DN41360\_c0\_g1 | CcBHLH53 | Helix-loop-helix DNA-binding domain | 1.02 | 0.78 | 4.90 |
| TRINITY\_DN45589\_c1\_g3 | CcBHLH54 | Helix-loop-helix DNA-binding domain | 22.97 | 25.08 | 7.88 |
| TRINITY\_DN35569\_c0\_g1 | CcBHLH55 | Helix-loop-helix DNA-binding domain | 3.41 | 104.09 | 10.62 |